



# scRNA-seq data processing

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 An abbreviated version of this protocol was published in Science Advances in Dec 2021


Methylome inheritance and enhancer dememorization reset an epigenetic gate safeguarding embryonic programs

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## Related files

 cell\_clusters.txt



 run\_seurat3.txt



**How to cite:** (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Xie, W. , Wu, X. and Zhang, H. (2022). scRNA-seq data processing. Bio-protocol Preprint. [bio-protocol.org/prep1626](https://bio-protocol.org/prep1626).
2. Wu, X., Zhang, H., Zhang, B., Zhang, Y., Wang, Q., Shen, W., Wu, X., Li, L., Xia, W., Nakamura, R., Liu, B., Liu, F., Takeda, H., Meng, A. and Xie, W.(2021). Methylome inheritance and enhancer dememorization reset an epigenetic gate safeguarding embryonic programs. Science Advances 7(52). DOI: [10.1126/sciadv.abl3858](https://doi.org/10.1126/sciadv.abl3858)

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